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Pubmed id 7751520

Title

7751520

Structural investigations of the major allergen PhI p I on the complementary

DNA and protein level.

Journal Article; Comparative Study;

Authors

Petersen A, Schramm G, Bufe A, Schlaak M, Becker WM

Affiliation

Division of Allergology, Forschungsinstitut Borstel, Germany.

Language

English

Journal

J. Allergy Clin. Immunol. (ISSN: 0091-6749) (ESSN: 1097-6825)

[1995 May; Volume: 95 (Issue: 5 Pt 1)] Page info: 987-94

Publication tvpe

Full text article

duis.

XML

E Abstract

Until now investigations of group I grass allergens have mainly been performed on ryegrass allergen (Lol p I). We studied this major allergen grass group with timothy grass pollen (PhI p I), a very common and important cause of type I allergy, to determine intraspecific and interspecific variations among different grass species. By immunoscreening a timothy grass pollen complementary DNA library we obtained three full-length clones. They revealed identical nucleotide sequences in the coding regions consisting of 262 amino acids, including a leader sequence of 23 amino acid residues. The comparison of our data with the amino acid sequences deduced from Lol p I and Hol 1 I clones showed sequence identities of greater than 85% and homologies of greater than 90%, indicating a high degree of sequence conservation. Despite the high degree of homology, amino acid differences were in immunodominant positions, which may be responsible for the differing immune response to group I allergens of different grass species.

⊞ Proteins

Uniprot Id Q4096Z

Name/Info

Polien allergen Phi pI precursor

Taxonomy Phleum pratense

⊞ Nucl. sequences EMBL id

PPRPHLP1X

Description P.pratense mRNA for pollen allergen PhlpI.

Seq Length 1152

E Referenced bv

⊞ Keywords (Mesh)

E Chemicals

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UniProtKB Entry

PIR View



Niceprot View | SRS View

UniProtKB Entry: Q40967

ENTRY NAME	Q40967 PHLPR New! View this entry in our Beta site			
ACCESSION NUMBER	Q40967			
Integrated into TrEMBL on	1996-11-01			
Sequence was last modified on	1999-01-01 (Sequence version 2)			
Annotations were last modified on	2007-09-11 (Entry version 42)			
NAME AND ORIGIN OF TH	E PROTEIN			
PROTEIN NAME	Pollen allergen Phl pl precursor			
GENE NAME	phlpI.p4			
SOURCE ORGANISM	Phleum pratense			
TAXONOMY ID	15957 [<u>NCBI</u> , <u>NEWT</u>]			
LINEAGE	Eukaryota; Viridiplantae; Streptophyta; Embryophyta; Tracheophyta; Spermatophyta Magnoliophyta; Liliopsida; Poales; Poaceae; BEP clade; Pooideae; Aveneae; Phleum			
PROTEIN EXISTENCE	Evidence at transcript level			
REFERENCES				
	Structural investigations of the major allergen Phl p I on the cDNA and protein level. 1995, J. Allergy Clin. Immunol., 95, 987-994. Position: NUCLEOTIDE SEQUENCE. PubMed: 7751520; Medline: 95270847.			
COMMENTS				
SIMILARITY	EA6; Belongs to the expansin family.			
SIMILARITY	EA29; Contains 1 expansin-like CBD domain.			
SIMILARITY	EA30; Contains 1 expansin-like EG45 domain.			
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DATABASE CROSS-REFER	ENCES			
EMBL	Z27090,CAA81613.1,mRNA. [GenBank, DDBJ]			
GENE3D	G3DSA:2.60.40.760,Expan_Lol_pI_C,1. G3DSA:2.40.40.40,Pollen_allergen,1.			
GO	GO:0005576, C:extracellular region, IEA: InterPro. GO:0019953, P:sexual reproduction, IEA: InterPro.			

	QuickGO	
GRAMENE	Q40967.	
HSSP	P43213, <u>1N10</u> .	•
INTERPRO	IPR007112,Expan_endogl. IPR007118,Expan_Lol_pI. IPR007117,Expan_Lol_pI_C. IPR005795,LolPI. IPR014734,Pollen_allergen. IPR005132,RlpA.	
PFAM	PF03330,DPBB_1,1. PF01357,Pollen_allerg_1,1.	
PIR	S38620, <u>S38620</u> .	
PRINTS	PR01225,EXPANSNFAMLY. PR00829,LOLP1ALLERGN.	
PROSITE	<u>PS50843</u> ,EXPANSIN_CBD,1. <u>PS50842</u> ,EXPANSIN_EG45,1.	
SMR	Q40967,25-262.	
UniRef	View cluster of proteins with at least 50% / 90% / 100% identity.	
KEYWORDS		
Secreted; Signal		

FEATURES				
Feature	Description	Begin Positio	n End Position	Length
SIGNAL PEPTIDE	POTENTIAL	1	23	23
CHAIN	pollen allergen Phl pI /FTId=PRO_5000147467	24	263	240

Feature sequence (Put the mouse on the feature above to see the sequence below):

IPKVPPGPNITATYGDKWLDAKSTWYGKPTAAGPKDNGGACGYKDVDKPPFSGM

SEQUENCE				
LENGTH	263 aa ·			
MOLECULAR WEIGHT	28203 Da			
CRC64 CHECKSUM	06D5EC94675335C6			
SEQUENCE	MASSSSVLLV VALFAVFLGS AHGIPKVPPG PNITATYGDK WLDAKSTWYG 50 KPTAAGPKDN GGACGYKDVD KPPFSGMTGC GNTPIFKSGR GCGSCFEIKC 100 TKPEACSGEP VVVHITDDNE EPIAAYHFDL SGIAFGSMAK KGDEQKLRSA 150 GEVEIQFRRV KCKYPEGTKV TFHVEKGSNP NYLALLVKFV AGDGDVVAVD 200 IKEKGKDKWI ALKESWGAIW RIDTPEVLKG PFTVRYTTEG GTKGEAKDVI 250 PEGWKADTAY ESK 263			

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BIBLIOGRAPHY	► <u>View Bibliography Information</u> ► <u>Submit Bibliography</u> PubMed: PMID: <u>7751520</u>	
PIRSF FAMILY	PIRSF019642 major pollen allergen/expansion	•
Biological Process GO:0019953: sexual reproduction [INTERPRO; evidence:IEA] Cellular Component GO:0005576: extracellular region [INTERPRO; evidence:IEA]		

STRUCTURE

PDB Similarity: <u>1BMW</u>(169-260,45.2%); <u>1WHO</u>(169-260,45.2%)

1BMW: SCOP CATH FSSP MMDB PDBsum 1WHO: SCOP CATH FSSP MMDB PDBsum

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General Information

Primary Accession # Z27090

Accession#

Z27090

SRS Entry ID

EMBL:Z27090 (formerly EMBL:PPRPHLP1X)

Molecule Type

linear mRNA

Sequence Length

1152

Entry Division

PLN (Plants)

Entry Data Class

STD (Standard)

Sequence Version

Z27090.1

Creation Date

09-NOV-1993

Modification Date

18-APR-2005

EMBL-SVA

Z27090

Description

Description

P.pratense mRNA for pollen allergen PhlpI.

Keywords

PhlpI; pollen allergen.;

Organism

Phleum pratense (timothy grass)

Organism

Classification

Eukaryota; Viridiplantae; Streptophyta; Embryophyta; Tracheophyta; Spermatophyta; Magnoliophyta; Liliopsida; Poales; Poaceae; BEP clade; Pooideae; Aveneae; Phleum.

References

1. Petersen, A.; Schramm, G.; Bufe, G.; Schlaak, M.; Becker, W.M.;

Structural investigations of the major allergen Phl p I on the complementary DNA and protein level

J. Allergy Clin. Immunol. 95(5 Pt 1):987-994 (1995)

DOI

10.1016/S0091-6749(95)70099-4

PubMed

7751520 (post) CiteXplore

Position

1-1149

2. Petersen, A.; Submitted (01-NOV-1993) to the EMBL/GenBank/DDBJ databases. Petersen A., Forschungsinstitut Borstel, Allergology, Parkallee 22, 23845 BORSTEL, Germany

Position

1-1149

Features

Key **Location Qualifier** Value

source 1..1152

organism Phleum pratense

strain agrostideae

isolate 24W28-7c

mol type mRNA

clone_lib cDNA/38/10/ZAP/6a

clone p4

cell_line Phleum pratense pollen/charge 24W28-7c

db xref taxon:15957

cds 9..800

gene phlpI.p4

product pollen allergen Phl pI

db_xref GOA:Q40967

db_xref HSSP:P43213

db_xref InterPro:IPR005132

db_xref InterPro:IPR005795

db_xref InterPro:IPR007112

db_xref InterPro:IPR007117

db xref InterPro:IPR007118

db_xref InterPro:IPR014734

db_xref <u>UniProtKB/TrEMBL:Q40967</u>

protein_id CAA81613.1

translation

>CAA81613.1

MASSSSVLLVVALFAVFLGSAHGIPKVPPGPNITATYGDK GGACGYKDVDKPPFSGMTGCGNTPIFKSGRGCGSCFEIKC EPIAAYHFDLSGIAFGSMAKKGDEQKLRSAGEVEIQFRRV NYLALLVKFVAGDGDVVAVDIKEKGKDKWIALKESWGAIW

sig_peptide 9..77

mat_peptide 78..797

gene phlpI.p4

gene phlpI.p4

product pollen allergen Phl pl

function unknown

3'utr 801..1152

Sequence

Characteristics

Length: 1152 BP, A Count:267, C Count:346, G Count:329, T

Count:210, Others Count:0

Sequence

>embl|Z27090|Z27090 P.pratense mRNA for pollen allergen PhlpI. aagacaagatggcttcttcctcatcggttctgctggtcgtggcgctgttcqccqtcttcc tgggctctgcgcatggcatccccaaggtccccccggcccgaacatcacggcgacctacg gcgacaagtggctggacgcgaagagcacctggtacggcaagccgacggccgccggtccca ${\tt aggacaacggcgcgcgtgcgggtacaaggacgtggacaagcccccgttcagcggcatga}$ ccggctgcggcaacacccccatcttcaagtccggccggggctgcggctcctgcttcgaga tcaagtgcaccaagcccgaggcctgctccggcgagcccgtggtggtccacatcaccgacg acaacgaggagcccatcgccgcgtaccacttcgacctctccggcatcgcgttcgggtcca tggccaagaagggcgacgagcagaagctgcgcagcgccggcgaggtggagatccagttcc gccgcgtcaagtgcaagtacccggagggcaccaaggtcaccttccacgtggagaaggggt ccaaccccaactacctggcgctgctggtgaagtttgtcgccggcgacggcgacgtqgtqq cggtggacatcaaggagaagggcaaggacaagtggatcgcgctcaaggagtcgtggggag ccatctggaggatcgacaccccggaggtgctcaagggccccttcaccgtccgctacacca ccgagggcggcaccaagggcgaggccaaggacgtcatccccgagggctggaaggccgaca ccgcctacgagtccaagtgaacaatccatccatcaatggatccttcaccacccctacgta $\verb|ccacttccggcaagttcaattttgactcaacaagatcaccaataagttaaccggtcggcc|\\$ acatctatataggccagcgcaagctcctagcatgttcatgctacagagaggggaaatgcg ccaacaaaaaatggtttgtcctccggcccacactcggaccgctttaattgtttatgtatc taagattgttcatgtttatgaacatatatattgtaatacataattttgtccataaaccat cgattatatcaa

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